



PCT10

RAW SEQUENCE LISTING DATE: 07/11/2002 PATENT APPLICATION: US/10/018,497A TIME: 13:30:44

Input Set : A:\EP.txt

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3 <110> APPLICANT: Lex M. Cowsert
            ISIS PHARMACEUTICALS, INC.
     6 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-13 EXPRESSION
     8 <130> FILE REFERENCE: RTSP-0046
--> 10 <140> CURRENT APPLICATION NUMBER: US/10/018,497A
--> 10 <141> CURRENT FILING DATE: 2001-12-12
   10 <150> PRIOR APPLICATION NUMBER: US 09/339,775
   11 <151> PRIOR FILING DATE: 1999-06-25
   13 <160> NUMBER OF SEQ ID NOS: 47
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   22 <222> LOCATION: (9)..(1073)
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   26
                   Met Gly Cys Thr Leu Ser Ala Glu Asp Lys Ala Ala Val
   27
                                                                                95
   29
         gag ega age aag atg ate gae ege aac tta egg gag gae ggg gaa aaa
         Glu Arg Ser Lys Met Ile Asp Arg Asn Leu Arg Glu Asp Gly Glu Lys
   30
   31
               15
                                   20
                                                                               143
   33
         gcg gcc aaa gaa gtg aag ctg ctg cta ctc ggt gct gga gaa tct ggt
   34
         Ala Ala Lys Glu Val Lys Leu Leu Leu Gly Ala Gly Glu Ser Gly
   35
                               35
                                                   40
   37
                                                                               191
         aaa agc acc att gtg aaa cag atg aaa atc att cat gag gat ggc tat
         Lys Ser Thr Ile Val Lys Gln Met Lys Ile Ile His Glu Asp Gly Tyr
   38
                                               55
   39
                           50
         tca gag gat gaa tgt aaa caa tat aaa gta gtt gtc tac agc aat act
                                                                               239
   41
   42
         Ser Glu Asp Glu Cys Lys Gln Tyr Lys Val Val Tyr Ser Asn Thr
   43
                                           70
                       65
                                                                               287
   45
         ata cag tee ate att gea ate ata aga gee atg gga egg eta aag att
   46
         Ile Gln Ser Ile Ile Ala Ile Ile Arg Ala Met Gly Arg Leu Lys Ile
   47
                                       85
   49
         gac ttt ggg gaa gct gcc agg gca gat gat gcc cgg caa tta ttt gtt
                                                                               335
   50
         Asp Phe Gly Glu Ala Ala Arg Ala Asp Asp Ala Arg Gln Leu Phe Val
   51
              95
                                  100
   53
         tta gct ggc agt gct gaa gaa gga gtc atg act cca gaa cta gca gga
                                                                               383
         Leu Ala Gly Ser Ala Glu Glu Gly Val Met Thr Pro Glu Leu Ala Gly
   54
   55
                              115
                                                  120
   57
         gtg att aaa egg tta tgg ega gat ggt ggg gta eaa get tge tte age
                                                                               431
   58
         Val Ile Lys Arg Leu Trp Arg Asp Gly Gly Val Gln Ala Cys Phe Ser
```

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5 0					130					135					140		
59	2022	taa	200	m22		car	ctc	aat	αat		act	tica	t.a.t.	tat		aat	479
61 62	Ara	Car	Ara	Glu	Tyr	Gln	Leu	Asn	Asp	Ser	Ala	Ser	Tvr	Tyr	Leu	Asn	
63	AIG	261	Arg	145	- y -	OIII	LCu		150	00-			- 1 -	155			
65	rat	cta	rat		ata	tcc	саσ	tet		tac	att	cca	act	cag	caa	qat	527
66	Acn	Leu	Aen	Δra	Tle	Ser	Gln	Ser	Asn	Tvr	Ile	Pro	Thr	Gln	Gln	Asp	
67	нэр	пец	160	nrg	110	DCI	0.1.1	165		-1-			170			•	
69	att	ctt		acq	аπа	ata	ааσ		aca	aac	att	gta	gaa	aca	cat	ttc	575
70	Val.	Len	Δra	Thr	Ara	Val	Lvs	Thr	Thr	Glv	Ile	Val	Glu	Thr	His	Phe	
71	Vul	175	**** 9	1111	*** 9	,	180			1		185					
73	acc		aaa	gac	cta	tac		aaσ	atq	ttt	gat	gta	ggt	ggc	caa	aga	623
74	Thr	Phe	Lvs	Asp	Leu	Tvr	Phe	Lvs	Met	Phe	Asp	Val	Gly	Gly	Gln	Arg	
75	190	1110				195		-1-			200		_	•-		205	
77		gaa	сда	aaa	ааσ		att	cac	tat	ttt	qaq	qqa	qtq	aca	gca	att	671
78	Ser	Glu	Ara	Lvs	LVS	Tro	Ile	His	Cvs	Phe	Glu	Gly	Val	Thr	Āla	Ile	
79	DCT	014	**** 9	₁	210				- 4	215		-			220		
81	atc	ttc	tat	ata		ctc	agt.	gat	tat	gac	ctt	qtt	ctq	gct	gag	gac	719
82	Tle	Phe	Cvs	Val	Ala	Leu	Ser	Asp	Tyr	Asp	Leu	Val	Leu	Ala	Glu	Asp	
83	110	1 -10	4 12	225				_	230	•				235			
85	σασ	σασ	atg		cga	atq	cat	qaa	aqc	atg	aaa	ctg	ttt	gac	agc	att	767
86	Glu	Glu	Met	Asn	Arq	Met	His	Ğlu	Ser	Met	Lys	Leu	Phe	Asp	Ser	Ile	
87			240					245					250				
89	tat	aat	aac	aaa	tgg	ttt	aca	gaa	act	tca	atc	att	ctc	ttc	ctt	aac	815
90	Cvs	Asn	Asn	Lys	Trp	Phe	Thr	Ğlu	Thr	Ser	Ile	Ile	Leu	Phe	Leu	Asn	
91	-	255		_			260					265					
93	aaq	aaa	gac	ctt	ttt	gag	gaa	aaa	ata	aag	agg	agt	ccg	tta	act	atc	863
94	Lys	Lys	Asp	Leu	Phe	Glu	Glu	Lys	Ile	Lys	Arg	Ser	Pro	Leu	Thr	Ile	
95	270	_				275					280					285	
97	tgt	tat	cca	gaa	tac	aca	ggt	tcc	aat	aca	tat	gaa	gag	gca	gct	gcc	911
98	Cys	Tyr	Pro	Glu	Tyr	Thr	Gly	Ser	Asn	Thr	Tyr	Glu	Glu	Ala	Ala	Ala	
99					290					295					300		
101	tai	t ati	t caa	a tgo	cag	g ttt	gaa	a gat	cto	g aad	c aga	a aga	a aaa	a gat	aco	c aag	959
102	Ту	r Ile	e Glı	ı Cys	s Glr	n Phe	e Glu	ı Asp) Le	ı Ası	n Ar	g Ar	g Lys			Lys	
103				305					310	_				315			1005
105	ga	g ato	c tai	act	cad	c tto	caco	c tgt	ge	c aca	a gad	cac	g aaq	g aat	gto	g cag	1007
106	Glı	ı Ile	е Ту	r Thi	His	s Phe	? Thi			a Thi	r Ası	o Thi			ı Val	l Gln	
107			320					325					330	-			1055
109	tt	t gti	t tti	: gat	gct	c gtt	aca	a gat	: gto	c ato	c ati	t aaa	a aad	c aac	tta	a aag	1055
110	Phe			e Asp) Ala	a Val								n Ası	ı Lei	ı Lys	
111		33) .				34		4			1112
113	gaa	a tgi	t gga	a cti	tat	t tga	a gaa	agcat	:gga	tgti	tagt	gaa a	agtta	actac	ca gu	gtggagt	g 1113
114			s Gl	y Lei	тул	r											
115	350																1173
117	tt	gaga	ccag	acad	cctti	ttg (etgto	ctcat	eg gg	ggca	gcta	c aa	gcat	gaac	ggga	accaggg	1233
119	aa	tggca	agca	gcat	gcag	gaa t	ctta	agcad	ct ci	ttta	gcaca	a ata	attti	cgta	ttag	ggaact	1293
121	tti	taati	tgac	atga	agato	get a	aagt	caga	ac at	ctgga	aacto	y ga	agaad	JUAE Laca	adag	gtgtgat	1353
123	to	gatc	gtca	agad	catca	act t	.gga1	ttct	t aa	ateti	caaca	a tg	cttai	Lgga	ayaı	tgtgaag	1413
125	tt	gaggt	tgct	gcat	tcta	aga a	CCCC	catat	g ta	ageti	Lacto	יייי כיייי		2000	+564	ctcttaa	1413
127	ac	cacca	agtg	gtto	catti	ctt a	aaggt	ודננו	LT Ca	atcaa	ayaga	a aga	aaLdi	はししし	Laci	caaattt	14/3

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154	<400> SEQUENCE: 3				
155	cagcactgcc agctaaaaca aat				23
158	<210> SEQ ID NO: 4				
159	<211> LENGTH: 22				
160	<212> TYPE: DNA				
161	<213> ORGANISM: Artificial Sequence				
163	<220> FEATURE:				
164	<223> OTHER INFORMATION: PCR Probe				
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167	agggcagatg atgcccggca at				22
170	<210> SEQ ID NO: 5				
171	<211> LENGTH: 19				
172	<212> TYPE: DNA				
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175	<220> FEATURE:				
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179					19
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185	<213> ORGANISM: Artificial Sequence				
187	<220> FEATURE:				
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			20
		getteee gtteteagee	20
		SEQ ID NO: 8	
		LENGTH: 18	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
		FEATURE:	
		OTHER INFORMATION: Antisense Oligonucleotide	
214		SEQUENCE: 8	1.0
215		ettgette getecaet	18
		SEQ ID NO: 9	
219	<211>	LENGTH: 18	
		TYPE: DNA	
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254	<210>	SEQ ID NO: 12	
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Input Set : A:\EP.txt

	<210> SEQ ID NO: 14	
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 VERIFICATION SUMMARY
 DATE: 07/11/2002

 PATENT APPLICATION: US/10/018,497A
 TIME: 13:30:45

Input Set : A:\EP.txt

Output Set: N:\CRF3\07112002\J018497A.raw

10 M:270 C: Current Application Number differs, Replaced Current Application No

10 M:271 C: Current Filing Date differs, Replaced Current Filing Date